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- <140> 10/808,052
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- Gly Lys Leu Gln Asp Ser Val Gly Tyr Arg Ile Ser Ser Asn Val Asp
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- Val Ala Leu Leu Trp Arg Asn Pro Asp Gly Asp Asp Gln Leu Ile
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- Gln Ile Thr Met Lys Asp Val Asn Val Glu Asn Val Asn Gln Gln Arg 85 90 95
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- Ser Ser Gly Thr Thr Asn Glu Val Asp Ile Ser Gly Asn Cys Lys Val 165 170 175
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- Ser Cys Lys Ile Ala Arg Ser Gly Phe Thr Thr Pro Asn Gln Val Leu 195 200 205
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- Ser Phe Val Ile Ala Val Leu Ala Glu Glu Thr His Asn Phe Gly Leu 225 230 235 240
- Asn Phe Leu Gln Thr Ile Lys Gly Lys Ile Val Ser Lys Gln Lys Leu 245 250 255
- Glu Leu Lys Thr Thr Glu Ala Gly Pro Arg Leu Met Ser Gly Lys Gln 260 265 270
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- Ile Val Gly Gln Val Phe Gln Ser His Cys Lys Gly Cys Pro Ser Leu 290 295 300

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Pro Arg Ser Ala Ser Thr Tyr Ser Leu Asp Ile Leu Tyr Ser Gly Ser 625 630 635 640

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Ala Gly Leu His Gly Ser Gln Val Val Ile Glu Ala Gln Gly Leu Glu 660 665 670

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Ser Gly Asp Pro Ile Ser Val Val Lys Gly Leu Ile Leu Leu Ile Asp 725 730 735

His Ser Gln Glu Leu Gln Leu Gln Ser Gly Leu Lys Ala Asn Ile Glu 740 745 750

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Val Ile Thr Thr Asp Ile Thr Val Asp Ser Ser Phe Val Lys Ala Gly 785 790 795 800

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<223> Wherein Xaa is any amino acid.

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Val Pro Gly Thr Ala Xaa Ser Arg Ser Ala Thr Arg Xaa Asn Cys Lys
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Xaa Glu Leu Glu Val Pro Gln Leu Cys Ser Phe Ile Leu Lys Xaa Ser 85 90 95

Gln Cys Thr Leu Lys Glu Val Tyr Gly Phe Asn Pro Glu Gly Lys Ala 100 105 110

Leu Leu Lys Lys Thr Lys Asn Ser Xaa Glu Xaa Ala Ala Ala Met Ser 115 120 125

Arg Xaa Glu Leu Lys Leu Ala Ile Pro Glu Gly Lys Gln Val Phe Leu 130 135 140

Tyr Pro Glu Lys Asp Glu Pro Thr Tyr Ile Leu Asn Ile Lys Arg Gly
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Ile Ile Ser Ala Leu Leu Val Pro Pro Glu Xaa Glu Glu Ala Lys Gln 165 170 175

Xaa Leu Phe Xaa Asp Thr Val Tyr Gly Asn Cys Ser Thr His Phe Thr 180 185 190

Val Lys Thr Arg Xaa Gly Asn Xaa Ala Thr Xaa Xaa Ser Thr Glu Arg 195 200 205

Asp Leu Gly Gln Cys Asp Arg Phe Lys Pro Ile Arg Thr Gly Ile Ser 210 220

Pro Xaa Ala Leu Ile Lys Gly Met Xaa Arg Pro Leu Ser Thr Leu Ile 225 230 235 240

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Val Arg Xaa Xaa Lys Pro Trp Ser Gln Gly Thr Phe Ser Asp Gln Gln 65 70 75 80

Trp Glu Thr Leu Gln His Ile Phe Arg Val Tyr Arg Ser Ser Phe Thr
85 90 95

Xaa Asp Xaa Lys Glu Xaa Ala Lys Xaa Xaa Arg Leu Ser Tyr Pro Leu 100 105 110

Glu Leu Gln Xaa Ser Ala Gly Cys Glu Xaa His Pro Gly Asn Ala Ser 115 120 125

Asn Asn Phe Phe His Val Ala Phe Gln Gly Lys Asp Ile Leu Ser Phe 130 135 140

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Leu Ala Xaa Gln Xaa Leu Asn Gln Asp Lys Trp Thr Xaa Glu Thr Xaa 165 170 175

Gln Trp Leu Leu Asn Gly Thr Cys Pro Gln Phe Val Ser Gly Leu Leu 180 185 190

Glu Ser Gly Lys Ser Glu Leu Lys Lys Gln Val Lys Pro Lys Xaa Trp 195 200 205

Leu Ser Arg Gly Pro Xaa Pro Xaa Pro Gly Arg Leu Leu Xaa Cys 210 215 220

His Val Ser Gly Xaa Tyr Pro Lys Pro Val Trp Val Lys Trp Xaa Xaa

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Xaa Asp Glu Thr Trp Tyr Leu Arg Ala Thr Leu Xaa Val Xaa Ala Gly 260 265 270

Glu Ala Xaa Gly Leu Ser Cys Arg Val Lys His Ser Ser Leu Xaa Gly 275 280 285

Gln Asp Ile Val Leu Tyr Trp Gly Gly Ser Tyr Thr Ser Met Gly Leu 290 295 300

Ile Ala Leu Ala Val Leu Ala Cys Leu Leu Phe Leu Leu Ile Val Gly 305 310 315 320

Phe Thr Ser Arg Phe Lys Arg Gln Thr Ser Tyr Gln Gly Val Leu 325 330 335

<210> 13

<211> 210

<212> PRT

<213> Homo sapiens

<220>

<221> VARIANT

<222> (1)..(210)

<223> Wherein Xaa is any amino acid.

<400> 13

Lys Cys Val Gln Ser Xaa Lys Pro Ser Leu Met Ile Gln Lys Ala Xaa 1 5 10 15

Xaa Gln Ala Leu Arg Lys Met Glu Pro Lys Asp Lys Asp Gln Glu Val 20 25 30

Leu Leu Gln Thr Phe Leu Asp Asp Ala Ser Pro Gly Asp Xaa Arg Xaa 35 40 45

Ala Ala Xaa Leu Met Xaa Xaa Arg Ser Pro Ser Gln Ala Asp Xaa Asn 50 55 60

Lys Ile Val Gln Xaa Leu Pro Trp Glu Gln Asn Glu Gln Val Lys Asn 65 70 75 80

Xaa Val Ala Xaa His Ile Ala Asn Xaa Leu Asn Ser Glu Glu Xaa Asp 85 90 95

Xaa Gln Asp Leu Lys Lys Leu Val Xaa Glu Ala Xaa Lys Glu Ser Gln
100 105 110

Leu Pro Thr Val Met Asp Phe Arg Lys Phe Ser Arg Asn Tyr Gln Leu 115 120 125

Tyr Lys Ser Val Xaa Leu Pro Ser Leu Asp Pro Xaa Ser Xaa Lys Ile

130 135 140

Glu Gly Asn Leu Xaa Phe Asp Pro Asn Asn Xaa Leu Pro Lys Glu Ser 145 150 155 160

Met Xaa Xaa Thr Thr Leu Thr Ala Phe Gly Phe Ala Ser Xaa Asp Xaa 165 170 175

Xaa Glu Ile Xaa Leu Glu Gly Lys Gly Phe Glu Pro Thr Leu Xaa Ala 180 185 190

Xaa Phe Gly Lys Gln Xaa Phe Phe Pro Xaa Ser Val Asn Lys Ala Leu 195 200 205

Tyr Trp 210

<210> 14

<211> 301

<212> PRT

<213> Homo sapiens

<220>

<221> VARIANT

<222> (1)..(301)

<223> Wherein Xaa is any amino acid.

<400> 14

Phe Ser Tyr Asn Asn Lys Tyr Gly Met Val Ala Gln Val Thr Gln Thr

Leu Lys Leu Glu Asp Thr Pro Lys Ile Asn Ser Arg Phe Phe Gly Glu 20 25 30

Gly Thr Xaa Lys Met Gly Leu Ala Xaa Glu Ser Thr Lys Ser Thr Ser
35 40 45

Pro Pro Lys Xaa Ala Glu Ala Val Xaa Xaa Leu Gln Glu Leu Lys
50 55 60

Lys Leu Thr Ile Ser Xaa Gln Xaa Ile Gln Arg Ala Xaa Leu Phe Asn 65 70 75 80

Xaa Xaa Val Thr Glu Leu Arg Gly Leu Ser Asp Glu Ala Val Thr Ser 85 90 95

Xaa Leu Pro Gln Leu Ile Glu Xaa Ser Ser Pro Xaa Xaa Leu Gln Ala 100 105 110

Leu Val Gln Cys Gly Xaa Pro Gln Cys Ser Thr His Ile Xaa Gln Xaa

Leu Lys Xaa Val His Ala Asn Pro Leu Leu Ile Asp Val Val Thr Tyr
130 . 135 140

Leu Val Ala Leu Xaa Pro Glu Pro Ser Ala Gln Gln Xaa Arg Glu Ile

Phe Asn Met Ala Arg Xaa Gln Arg Ser Arg Ala Thr Leu Tyr Ala Leu 165 170 175

Ser His Ala Val Asn Asn Tyr His Lys Xaa Asn Pro Xaa Gly Thr Gln 180 185 190

Glu Leu Xaa Asp Ile Ala Asn Xaa Leu Met Glu Gln Ile Gln Asp Asp 195 200 205

Cys Xaa Gly Asp Glu Asp Tyr Thr Tyr Leu Xaa Leu Arg Xaa Ile Gly 210 215 220

Asn Met Gly Gln Thr Met Glu Gln Leu Thr Pro Glu Leu Lys Ser Xaa 225 230 235 240

Ile Leu Lys Cys Val Gln Ser Thr Lys Pro Ser Xaa Xaa Ile Gln Lys 245 250 255

Ala Ala Ile Gln Xaa Leu Arg Lys Met Glu Pro Lys Asp Lys Asp Gln 260 265 270

Xaa Xaa Leu Leu Gln Thr Phe Leu Asp Asp Ala Ser Pro Gly Asp Lys 275 280 285

Arg Leu Ala Ala Tyr Leu Met Leu Xaa Arg Ser Pro Ser 290 295 300

<210> 15

<211> 335

<212> PRT

<213> Homo sapiens

<220>

<221> VARIANT

<222> (1)..(335)

<223> Wherein Xaa is any amino acid.

<400> 15

Met Gly Cys Leu Leu Phe Leu Leu Trp Ala Leu Leu Gln Ala Trp 1 5 10 15

Gly Ser Ala Glu Val Pro Gln Arg Leu Phe Pro Leu Arg Cys Leu Gln 20 25 30

Ile Ser Ser Phe Ala Asn Ser Ser Trp Thr Arg Thr Asp Gly Leu Ala
35 40 45

Trp Leu Gly Glu Leu Gln Thr His Xaa Trp Ser Asn Asp Ser Asp Thr
50 55 60

Val Arg Xaa Xaa Lys Pro Trp Ser Gln Gly Thr Phe Ser Asp Gln Gln 65 70 75 80

Trp Glu Thr Leu Gln His Ile Phe Arg Val Tyr Arg Ser Ser Phe Thr

85	90	95
05	70	

Xaa	Asp	Xaa	Lys 100	Glu	Xaa	Ala	Lys	Xaa 105	Xaa	Arg	Leu	Ser	Tyr 110	Pro	Leu
Glu	Leu	Gln 115	Xaa	Ser	Ala	Gly	Cys 120	Glu	Xaa	His	Pro	Gly 125	Asn	Ala	Ser
Asn	Asn 130	Phe	Phe	His	Val	Ala 135	Phe	Gln	Gly	Lys	Asp 140	Ile	Leu	Ser	Phe
Gln 145	Gly	Thr	Ser	Xaa	Glu 150	Pro	Xaa	Gln	Glu	Ala 155	Pro	Xaa	Trp	Val	Asn 160
Leu	Ala	Xaa	Gln	Xaa 165	Leu	Asn	Gln	Asp	Lys 170	Trp	Thr	Xaa	Glu	Thr 175	Xaa
Gln	Trp	Leu	Leu 180	Asn	Gly	Thr	Cys	Pro 185	Gln	Phe	Val	Ser	Gly 190	Leu	Leu
Glu	Ser	Gly 195	Lys	Ser	Glu	Leu	Lys 200	Lys	Gln	Val	Lys	Pro 205	Lys	Xaa	Trp
Leu	Ser 210	Arg	Gly	Pro	Xaa	Pro 215	Xaa	Pro	Gly	Arg	Leu 220	Leu	Leu	Xaa	Cys
His 225	Val	Ser	Gly	Xaa	Tyr 230	Pro	Lys	Pro	Val	Trp 235	Val	Lys	Trp	Xaa	Xaa 240
Gly	Glu	Gln	Glu	Gln 245	Gln	Gly	Thr	Gln	Pro 250	Xaa	Asp	Xaa	Xaa	Pro 255	Asn
Xaa	Asp	Glu	Thr 260	Trp	Tyr	Leu	Arg	Ala 265	Thr	Leu	Xaa	Val	Xaa 270	Ala	Gly
Glu	Ala	Xaa 275	Gly	Leu	Ser	Cys	Arg 280	Val	Lys	His	Ser	Ser 285	Leu	Xaa	Gly

Gln Asp Ile Val Leu Tyr Trp Gly Gly Ser Tyr Thr Ser Met Gly Leu

Ile Ala Leu Ala Val Leu Ala Cys Leu Leu Phe Leu Leu Ile Val Gly

Phe Thr Ser Arg Phe Lys Arg Gln Thr Ser Tyr Gln Gly Val Leu

<210> 16

305

<211> 335

290

<212> PRT

<213> Homo sapiens

<220>

<221> VARIANT

<222> (1)..(335)

<223> Wherein Xaa is any amino acid.

325

310

330

320

<400> 16 Met Gly Cys Leu Leu Phe Leu Leu Leu Trp Ala Leu Leu Gln Ala Trp 10 Gly Ser Ala Glu Val Pro Gln Arg Leu Phe Pro Leu Arg Cys Leu Gln Ile Ser Ser Phe Ala Asn Ser Ser Trp Thr Xaa Thr Asp Gly Leu Ala Xaa Leu Gly Glu Leu Gln Thr His Ser Trp Ser Xaa Asp Ser Asp Thr Xaa Xaa Xaa Leu Lys Pro Trp Ser Gln Gly Thr Phe Ser Xaa Gln Xaa 75 Trp Glu Thr Leu Xaa His Ile Phe Xaa Xaa Tyr Arg Ser Ser Phe Thr Arg Asp Val Lys Glu Phe Ala Lys Xaa Leu Arg Leu Ser Tyr Pro Xaa Glu Leu Gln Xaa Xaa Ala Gly Cys Glu Val His Pro Gly Xaa Ala Ser 120 Asn Asn Phe Phe His Xaa Ala Xaa Gln Gly Xaa Asp Ile Leu Ser Phe 130 135 Gln Gly Thr Ser Trp Glu Pro Thr Gln Glu Ala Pro Xaa Trp Val Asn 150 155 Leu Ala Ile Gln Xaa Leu Asn Gln Asp Lys Trp Thr Arg Xaa Thr Val Gln Trp Leu Leu Asn Gly Thr Cys Pro Gln Phe Val Ser Gly Leu Leu 180 185 Glu Xaa Gly Lys Xaa Glu Leu Lys Lys Gln Xaa Lys Pro Lys Ala Xaa 200 Leu Ser Arg Gly Pro Ser Pro Gly Pro Gly Arg Leu Leu Val Cys 210 215 His Val Xaa Gly Phe Tyr Pro Lys Pro Val Trp Xaa Lys Trp Xaa Arg 235

Gly Glu Gln Gln Gln Gln Gln Gly Thr Gln Pro Gly Asp Ile Leu Pro Asn 245 Asp Asp Gly Asp Ile Leu Pro Asn 255 Asn Xaa Asp Gly Thr 260 Trp Tyr Leu Arg Ala Thr Leu Asp Xaa Xaa Ala Gly Glu Ala Ala Gly 275 Clu Xaa Cys Arg Val Lys His Ser Ser Leu Glu Gly

Gln Xaa Xaa Xaa Leu Tyr Trp Gly Gly Ser Tyr Thr Ser Met Gly Leu

	290					295					300					
Ile 305	Ala	Leu	Ala	Val	Leu 310	Ala	Cys	Leu	Xaa	Phe 315	Leu	Leu	Ile	Val	Gly 320	
Phe	Thr	Ser	Arg	Phe 325	Lys	Arg	Gln	Thr	Ser 330	Tyr	Gln	Gly	Val	Leu 335		